RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	/o/535,4/4A
Source:	IFWR
Date Processed by STIC:	8/8/06
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ENTERED



IFWP

RAW SEQUENCE LISTING DATE: 08/08/2006
PATENT APPLICATION: US/10/535,414A TIME: 09:22:32

Input Set : A:\Q87920.ST25.txt

3 <110> APPLICANT: SHARMA, Praveen 4 SAHNI, Narinder Singh

```
LONNEBORG, Anders
      7 <120> TITLE OF INVENTION: PRODUCT AND METHOD
      9 <130> FILE REFERENCE: Q87920
     11 <140> CURRENT APPLICATION NUMBER: US 10/535,414A
     12 <141> CURRENT FILING DATE: 2005-05-19
     14 <150> PRIOR APPLICATION NUMBER: PCT/GB03/05102
     15 <151> PRIOR FILING DATE: 2003-11-21
     17 <160> NUMBER OF SEQ ID NOS: 501
     19 <170> SOFTWARE: PatentIn version 3.3
     21 <210> SEQ ID NO: 1
     22 <211> LENGTH: 405
     23 <212> TYPE: DNA
     24 <213> ORGANISM: Homo sapiens
     26 <400> SEQUENCE: 1
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     27 ggatectgtg geceacagag etgececage agaegeteeg eeceaceegg tgatggagee
     29 ccggggggac aatcgtgcct ggggaggagc agggtacagc ccattccccc agccctggct
     31 gacctggcct agcagtttgg ccctgctggc cttagcaggg agacagggga gcaaagaacg
                                                                              180
     33 ccaageegga ggeeegagge cageeggeet etegagagee agageageag ttgaatgtaa
                                                                              240
     35 tgctggggac aggcatgctg ccgccagtag ggcggggacc cggacagcca ggtgactacc
                                                                              300
                                                                              360
     37 agtcctgggg acacactcac cataaacaca tccccaggca ggacagatcg gggaaggggt
     39 gtgtaccagg ctatgatttc tcttgcatta aaatgtatta ttatt
                                                                              405
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     43 <211> LENGTH: 550
     44 <212> TYPE: DNA
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     49 <221> NAME/KEY: misc feature
     50 <222> LOCATION: (61)..(61)
     51 <223> OTHER INFORMATION: n is a, c, g, or t
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     54 <221> NAME/KEY: misc feature
     55 <222> LOCATION: (464)..(464)
     56 <223> OTHER INFORMATION: n is a, c, g, or t
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                                                                              60
W--> 61 naccatcatc aacaccttcc accaatactc tgtgaagctg gggcacccag acaccctgaa
                                                                              120
     63 ccaqqqqqaa ttcaaaqagc tqgtqcgaaa agatctgcaa aattttctca agaaggagaa
                                                                              180
     65 taagaatgaa aaggtcatag aacacatcat ggaggacctg gacacaaatg cagacaagca
                                                                              240
    67 getgagette gaggagttea teatgetgat ggegaggeta acetgggeet eecaegagaa
                                                                              300
                                                                              360
     69 qatqcacqag ggtgacgagg gccctggcca ccaccataag ccaggcctcg gggagggcac
     71 cccctaagac cacagtggcc aagatcacag tggccacggc cacqqccaca gtcatggtgg
                                                                              420
```

RAW SEQUENCE LISTING DATE: 08/08/2006
PATENT APPLICATION: US/10/535,414A TIME: 09:22:32

Input Set : A:\Q87920.ST25.txt

	73 ccacggccac agccactaat caggaggcca ggccaccctg cctntaccca accagggccc 75 cggggcctgt tatgtcaaac tgtcttggct gtggggctag gggctggggc caaataaagt	480 540
	77 ctctttctcc	550
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	82 <212> TYPE: DNA	
	83 <213> ORGANISM: Homo sapiens	
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	86 acgaagacag acatctgtgg aatgattcac atcctctcaa gttaggagga tggaggcctg	60
	88 cttcattaag aagctggggg tagggtgggg gtggggagaa cacttaacaa catggggacc	120
	90 agtcagggga atccccttat ttctgttttg catatgagga accctagagc agccaggtga	180
	92 ggctctctag tttaataaaa atcatggaaa gactcttaat gcagactctt cttaagtgtt	240
	94 aatagggatt ttttcagctt attttggttg cagtttccaa tttttaaaaa tgttgaggta	300
	96 atctttccca ccttcccaaa cctaattctt gtagatgcat tagtgttgaa ccaatgcttt	360
	98 ctcatgtctc aattctttgt atatgcattc ttttcagatg tattaaacaa acaaaaaccc	420
	100 ttc	423
	103 <210> SEQ ID NO: 4	
	104 <211> LENGTH: 286	
	105 <212> TYPE: DNA	
	106 <213> ORGANISM: Homo sapiens	
	109 <220> FEATURE:	
	110 <221> NAME/KEY: misc_feature	
	111 <222> LOCATION: (64)(65)	
	112 <223> OTHER INFORMATION: n is a, c, g, or t	
	114 <400> SEQUENCE: 4	
	115 ccggtaatag aatagaaaag ggagagtgtc ttcatgcaat gtggcatcct ggattgggtc	60
W>	117 tcgnnacaaa aacaggacat tagtgggaaa attggaaatc tgaaaaaagt ctgaatttta	120
	119 gttaatatac caatttcagt ctcttggttt tgacagatgt accatggtga tgtaagatgt	180
	121 tgaccttggg gtaggctggg tgaagggtat acaggaactc tttgtactat ctctgcaact	240
	123 tctctgtaaa tctagtatca ttccaaaata aaagtttatt taattt	286
	126 <210> SEQ ID NO: 5	
	127 <211> LENGTH: 545	
	128 <212> TYPE: DNA	
	129 <213> ORGANISM: Homo sapiens	
	131 <400> SEQUENCE: 5	
	132 gtggaagtga catcgtcttt aaaccctgcg tggcaatccc tgacgcaccg ccgtgatgcc	60
	134 cagggaagac agggcgacct ggaagtccaa ctacttcctt aagatcatcc aactattgga	120
	136 tgattatccg aaatgtttca ttgtgggagc agacaatgtg ggctccaagc agatgcagca	180
	138 gatccgcatg tcccttcgcg ggaaggctgt ggtgctgatg ggcaagaaca ccatgatgcg	240
	140 caaggccatc cgagggcacc tggaaaacaa cccagctctg gagaaactgc tgcctcatat	300
	142 ccgggggaat gtgggctttg tgttcaccaa ggaggacctc actgagatca gggacatgtt	360
	144 gctggccaat aaggtgccag ctgctgcccg tgctggtgcc attgccccat gtgaagtcac	420
	146 tgtgccagcc cagaacactg gtctcgggcc cgagaagacc tcctttttcc aggctttagg	480
	148 tatcaccact aaaatctcca ggggcaccat tgaaatcctg agtgatgtgc actgatcaag	540
	150 actgg	545
	153 <210> SEQ ID NO: 6	
	154 <211> LENGTH: 591	
	155 <212> TYPE: DNA	
	156 <213> ORGANISM: Homo sapiens	

RAW SEQUENCE LISTING DATE: 08/08/2006 PATENT APPLICATION: US/10/535,414A TIME: 09:22:32

Input Set : A:\Q87920.ST25.txt

	159	<220> FEATURE:	
	160	<221> NAME/KEY: misc feature /	
		<222> LOCATION: (85)(85)	
		<223> OTHER INFORMATION: n is a, c, g, or t	
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		<222> LOCATION: (485)(485)	
		<223> OTHER INFORMATION: n is a, c, g, or t	
		<400> SEQUENCE: 6	
	170	cagegeaggg gettetgetg agggggeagg eggagettga ggaaacegea gataagtttt	60
W>			120
	174	aagatattgc ttagcgttaa gtttttaacg taattttaat agcttaagat tttaagagaa	180
			240
	178	acggaggttg agatgaagct tcttcatgga gtaaaaaatg tatttaaaag aaaattgaga	300
	180	gaaaggacta cagagccccg aattaatacc aatagaaggg caatgctttt agattaaaat	360
	182	gaaggtgact taaacagctt aaagtttagt ttaaaagttg taggtgatta aaataatttg	420
	184	aaggcgatct tttaaaaaaga gattaaaccg aaggtgatta aaagaccttg aaatccatga	480
	186	cgcanggaga attgcgcatt taaagcctag ttacgcattt actaaacgca gacgaaaatg	540
			591
	191	<210> SEQ ID NO: 7	
	192	<211> LENGTH: 297	
	193	<212> TYPE: DNA	
		<213> ORGANISM: Homo sapiens	
	196	<400> SEQUENCE: 7	
		ctcaaaggag aaaaaaacc ttgtaaaaaa agcaaaaatg acaacagaaa aacaatctta	60
		3 3 33	120
		-33-3 333 33 3	180
			240
			297
		<210> SEQ ID NO: 8	
		<211> LENGTH: 282	
		<212> TYPE: DNA	
		<213> ORGANISM: Homo sapiens	
		<400> SEQUENCE: 8	
		agtagagacg gggtttcact gtgttagcca ggatggtctc gatctcctga cctcgtgatc	60
			120
			180 240
		2.	282
			202
		<210> SEQ ID NO: 9	
		<211> LENGTH: 619 <212> TYPE: DNA	
		<212> TIPE: DNA <213> ORGANISM: Homo sapiens	
		<213> ORGANISM: HOMO SAPIEMS <400> SEQUENCE: 9	
		tccttggttt cgatttgtgg caacaatcca gtctttttgt ttttttcagg gataccatat	60
			120
			180
			240
			300
	433	aactaattii gagtigitty gataaaaatg tittagtiga tigtagtgig tattgagaga	500

DATE: 08/08/2006

TIME: 09:22:32

Input Set : A:\Q87920.ST25.txt Output Set: N:\CRF4\08082006\J535414A.raw 241 cactgccagt aaacaaactc tcttggtagg tggaaatccc ctagaagtta cagaaaattg 243 ggaggaggtg aacttaatta aataacttga attgtttaga catattcaga gcttcttatg 420 245 accttgaaga aatcacccaa cttcaaaaga cctcggtttc ttcatttgta aaattaggga 480 247 gtttgactag atgtgtaaat ctagttgtta gttaacttct aagatgtaaa aaccctcttg 540 249 tttaacaaaa acctacaaga tcaagttgct tatctgaaat ctttatgaat caacactagt 600 619 251 cactaagtct agctcgacc 254 <210> SEQ ID NO: 10 255 <211> LENGTH: 536 256 <212> TYPE: DNA 257 <213> ORGANISM: Homo sapiens 260 <220> FEATURE: 261 <221> NAME/KEY: misc feature 262 <222> LOCATION: (513)..(513) 263 <223> OTHER INFORMATION: n is a, c, g, or t 265 <220> FEATURE: 266 <221> NAME/KEY: misc feature 267 <222> LOCATION: (520)..(520) 268 <223> OTHER INFORMATION: n is a, c, g, or t 270 <220> FEATURE: 271 <221> NAME/KEY: misc feature 272 <222> LOCATION: (529)..(529) 273 <223> OTHER INFORMATION: n is a, c, g, or t 275 <400> SEQUENCE: 10 276 cttttcctcc cgctgtcccc cacggaggg actgctctcc cccgctgcat cctttctgtg 278 aggtacctta cccacctcag cacctgagag ggtgaaatag aattctaacc tcgacattcg 120 280 ggaagtgttt ttgagaagtc tcggtcggta agggaagtct tccaagtccg tgcagcacta 180 282 acgtattggc acctgcctcc tettcggcca cccccagat gaggcagctg tgactgtgtc 240 284 aagggaagee aegaetetga eeatagtett eteteagett eeactgeegt eteeacagga 300 286 aacccagaag ttctgtgaac aagtccatgc tgccatcaag gcatttattg cagtgtacta 360 288 tttgcttcca aaggatcagg ccctgagaac aatgacctta tttcctacaa cagtgtctgg 420 290 gttgcgtgcc agcagatgcc tcagatacca agagataaca aagctgcagc tcttttgatg 480 W--> 292 ctgaccaaga atgtggattt tgtgaaggat gcncatgaan aaatggacna gctgtg 536 295 <210> SEQ ID NO: 11 296 <211> LENGTH: 373 297 <212> TYPE: DNA 298 <213> ORGANISM: Homo sapiens 301 <220> FEATURE: 302 <221> NAME/KEY: misc feature 303 <222> LOCATION: (27)..(27) 304 <223> OTHER INFORMATION: n is a, c, g, or t 306 <220> FEATURE: 307 <221> NAME/KEY: misc_feature 308 <222> LOCATION: (235)..(235) 309 <223> OTHER INFORMATION: n is a, c, g, or t 311 <220> FEATURE: 312 <221> NAME/KEY: misc_feature 313 <222> LOCATION: (248)..(248) 314 <223> OTHER INFORMATION: n is a, c, g, or t 316 <220> FEATURE:

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/535,414A

RAW SEQUENCE LISTING DATE: 08/08/2006
PATENT APPLICATION: US/10/535,414A TIME: 09:22:32

Input Set : A:\Q87920.ST25.txt

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    318 <222> LOCATION: (329)..(329)
    319 <223> OTHER INFORMATION: n is a, c, g, or t
    321 <220> FEATURE:
    322 <221> NAME/KEY: misc_feature
    323 <222> LOCATION: (335)..(335)
    324 <223> OTHER INFORMATION: n is a, c, g, or t
    326 <220> FEATURE:
    327 <221> NAME/KEY: misc_feature
     328 <222> LOCATION: (359)..(359)
     329 <223> OTHER INFORMATION: n is a, c, g, or t
    331 <220> FEATURE:
    332 <221> NAME/KEY: misc feature
     333 <222> LOCATION: (372)..(372)
     334 <223> OTHER INFORMATION: n is a, c, g, or t
    336 <400> SEQUENCE: 11
W--> 337 aagtgggtet tgccatecet gaactgnaat catecetaac atatteatae etgtttteat
                                                                              60
     339 tttaaaaqtt qqqtcaqttt ttttattaqt acatqtattt ctatcctact gatttatttg
                                                                             120
     341 ctatatcatc taatttagtt tgaatattcc ataatttact taattagtcc tgtatggaga
                                                                             180
     343 cctagctctt ctcagtgtct actattataa acaatgctac agtgaatatt ggtgnataaa
                                                                             240
     345 tecataenea eeaegtaeat atettaagtt etggaagaga tattgetaaa eeagaagata
                                                                             300
     347 acctqcattt aaaatttqac tqctaqgqnc agggncacat ttaattaaat tagaacaang
                                                                             360
                                                                             373
     349 aatqcataat qnc
     352 <210> SEQ ID NO: 12
    353 <211> LENGTH: 796
     354 <212> TYPE: DNA
     355 <213> ORGANISM: Homo sapiens
     358 <220> FEATURE:
     359 <221> NAME/KEY: misc_feature
     360 <222> LOCATION: (601)..(601)
     361 <223> OTHER INFORMATION: n is a, c, g, or t
     363 <400> SEQUENCE: 12
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     366 taggcaatct gaaggaaacc agaaaatgtg aatttctctt ccctcaaaaa gctatactga
                                                                             120
     368 agtagtattt aatattcaag tacttgtaaa tttgcagaac agtacttttt aatttgaccc
                                                                             180
     370 atgaattota tttaaatttg toacttaata tttagocaag aagcaaacca totaaaaaga
                                                                             240
     372 tttctqqttt atttctccaa ctcctaataa atagggtcac atatttttta actttttct
                                                                             300
     374 aatttqaaaa qtaatacaqq catatqqtat tttaaaaaatg aaacaacaca aagggatatg
                                                                             360
     376 ttttgaaaag tggtcttgcc atccctgaac tgtaatcatc cctaacatat tcatacctgt
     378 tttcatttta aaagttgggt cagttttttt attagtacat gtatttctat cctactgatt
                                                                             480
                                                                             540
     380 tatttqctat atcatctaat ttaqtttqaa tattccataa tttacttaat taqtcctgta
     382 tggagaccta gctcttctca gtgtctacta ttataaacaa tgctacagtg aatattggtg
                                                                             600
W--> 384 nataaatcct acacaccacg taacatatct taagttcctg gaagagatat tgctaaacca
                                                                             660
     720
                                                                             780
     388 gaacaaggaa tgcataatgt cttcgatagc aatctattca aggtgcaccg tggtcacaaa
     390 ggaaagcaaa actgtc
                                                                             796
     393 <210> SEQ ID NO: 13
     394 <211> LENGTH: 564
     395 <212> TYPE: DNA
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/08/2006
PATENT APPLICATION: US/10/535,414A TIME: 09:22:33

Input Set : A:\Q87920.ST25.txt

Output Set: N:\CRF4\08082006\J535414A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

```
Seq#:2; N Pos. 61,464
Seq#:4; N Pos. 64,65-
Seq#:6; N Pos. -85,485
Seq#:10; N Pos. 513,520,529
Seq#:11; N Pos. 27,235,248,329,335,359,372
Seq#:12: N Pos. 507
Seq#:12; N Pos. 601/
Seq#:13; N Pos. 6,26,55,73,99,100,180,228,486
Seq#:15; N Pos. 149,177,463
Seq#:20; N Pos. 557
Seq#:22; N Pos. 199
Seq#:24; N Pos. 149,177,463
Seg#:25; N Pos. 388,506,672
Seq#:26; N Pos. 524
Seq#:28; N Pos. 213,306
Seq#:31; N Pos. 448,617,663,677,706,710
Seq#:32; N Pos. 53,89
Seq#:34; N Pos. 5,8,21,25,65,79,80,103,143,481
Seq#:35; N Pos. 6
Seq#:36; N Pos. 525
Seq#:42; N Pos. 119,121,157,205,214,216,229,252,327,329,354,361
Seq#:45; N Pos. 304
Seq#:47; N Pos. 7
Seq#:55; N Pos. 462
Seq#:57; N Pos. 571
Seq#:63; N Pos. 441
Seq#:65; N Pos. 270,280,310,359
Seq#:66; N Pos. 195,201
Seq#:67; N Pos. 606
Seq#:68; N Pos. 92,642
Seq#:69; N Pos. 44
Seq#:70; N Pos. 684,707,709,713,723,734,746,748
Seq#:71; N Pos. 8
Seq#:73; N Pos. 280,342,500
Seq#:74; N Pos. 552
Seq#:76; N Pos. 461,483,497
Seq#:77; N Pos. 567,575
Seq#:84; N Pos. 223,280,297,302,337
Seq#:85; N Pos. 9,29,49,94,105,152,169
Seq#:88; N Pos. 529,542,559
Seq#:93; N Pos. 70,75,88,102,114,127,129
Seq#:95; N Pos. 650,685
Seq#:96; N Pos. 204,255
Seq#:97; N Pos. 248,519
Seq#:98; N Pos. 425
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/08/2006
PATENT APPLICATION: US/10/535,414A TIME: 09:22:33

Input Set : A:\Q87920.ST25.txt

Output Set: N:\CRF4\08082006\J535414A.raw

Seq#:99; N Pos. 552

Seq#:101; N Pos. 565,581

Seq#:102; N Pos. 86,168,230,241,257,337,339
Seq#:103; N Pos. 20,22,47,546,562,611,645

Seq#:107; N Pos. 708
Seq#:109; N Pos. 600
Seq#:110; N Pos. 616,635

VERIFICATION SUMMARYDATE: 08/08/2006PATENT APPLICATION: US/10/535,414ATIME: 09:22:33

Input Set : A:\Q87920.ST25.txt

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L:61 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:60
M:341 Repeated in SeqNo=2
L:117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:60
L:172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:60
M:341 Repeated in SeqNo=6
L:292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:480
L:337 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
M:341 Repeated in SeqNo=11
L:384 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:600
L:440 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
M:341 Repeated in SeqNo=13
L:502 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:120
M:341 Repeated in SeqNo=15
L:641 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:540
L:697 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:180
L:771 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:120
M:341 Repeated in SeqNo=24
L:822 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:360
M:341 Repeated in SeqNo=25
L:863 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:480
L:914 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:180
M:341 Repeated in SeqNo=28
L:1042 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:420
M:341 Repeated in SeqNo=31
L:1072 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
M:341 Repeated in SeqNo=32
L:1168 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0
M:341 Repeated in SeqNo=34
L:1203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0
L:1254 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:480
L:1411 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:60
M:341 Repeated in SeqNo=42
L:1496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:300
L:1536 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:0
L:1764 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:420
L:1826 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57 after pos.:540
L:1988 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63 after pos.:420
L:2055 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65 after pos.:240
M:341 Repeated in SeqNo=65
L:2085 \ M:341 \ W: (46) "n" or "Xaa" used, for SEQ ID#:66 after pos.:180
L:2120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:600
L:2142 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68 after pos.:60
M:341 Repeated in SeqNo=68
L:2175 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69 after pos.:0
L:2267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 after pos.:660
M:341 Repeated in SeqNo=70
L:2284 \ M:341 \ W: (46) "n" or "Xaa" used, for SEQ ID#:71 after pos.:0
L:2342 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73 after pos.:240
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VERIFICATION SUMMARY

DATE: 08/08/2006 TIME: 09:22:33

PATENT APPLICATION: US/10/535,414A

Input Set : A:\Q87920.ST25.txt
Output Set: N:\CRF4\08082006\J535414A.raw

M:341 Repeated in SeqNo=73

. . . .

L:2383 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74 after pos.:540